

D. Kruse

Re-run

#4



RAW SEQUENCE LISTING

DATE: 01/14/2003

PATENT APPLICATION: US/09/954,773A

TIME: 14:52:26

Input Set : N:\paola\US09954773A.raw

Output Set: N:\CRF4\01142003\I954773A.raw

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1 <110> APPLICANT: Southern Illinois University
2     Lightfoot, David A
3     Gibson, Paul T
4     Meksem, Khalid
5 <120> TITLE OF INVENTION: SOYBEAN SUDDEN DEATH SYNDROME RESISTANT SOYBEANS, SOYBEAN
CYST NEMATODE
6     RESISTANT SOYBEANS AND METHODS OF BREEDING AND IDENTIFYING RESISTANT PLANTS
7 <130> FILE REFERENCE: 1268/2/2
C--> 8 <140> CURRENT APPLICATION NUMBER: US/09/954,773A
9 <141> CURRENT FILING DATE: 2002-01-22
10 <150> PRIOR APPLICATION NUMBER: 09/007,119
11 <151> PRIOR FILING DATE: 1998-01-14
12 <160> NUMBER OF SEQ ID NOS: 20
13 <170> SOFTWARE: PatentIn version 3.1
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17 <212> TYPE: DNA
18 <213> ORGANISM: Glycine max
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21 <222> LOCATION: (1)..(527)
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26     ttttattaaa taataaaaga gtaattacat aaaaatatgt tcgattacat taaaccccaa      180
27     caaaggatga atttagcttc tcatgacctt ggggaaaatc aaacttgatg aacaagaaga      240
28     tgaagaagaa tccttaagga taaacactgc ctagctccaa tgtgctctct agtattttat      300
29     ctttcaaaaa tccccaagaa cccctaattt tcagtaagaa gcccatTTTC aatcagaagc      360
30     ccattttcaa tcaagaagcc cattttcaat cagaagccca ttttcaatca gaagcccatt      420
31     ttcaatcaga agcccatttt caatcagaag cccattttat aattgtattc ccaaaacttg      480
32     agattcttga acgtaaatta ttagtaaatt gtaatcacct ctgtaaa                    527
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35 <211> LENGTH: 815
36 <212> TYPE: DNA
37 <213> ORGANISM: Glycine max
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39 <221> NAME/KEY: misc_feature
40 <222> LOCATION: (1)..(815)
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44     gttgggagct ctcccatatg gtcgacctgc aggcggccgc actagtattt cagaagccca      120
45     aaggaacag caataagtaa tcccttgttt ataagatccc agaacttcca gtttatttaa      180

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46      tgaaaatgca ataacatcgg ctagtttcac aagtaatata caaatcggaa catcacattg      240
47      actacaatat atagtacata aattaacact aagaaacctc cttgatttga tattatgcat      300
48      ttacctatgt tgttccacaa gaatatactc aaatgacttt gccttgattt aaattatcac      360
W--> 49      gatgtaacac aaacaaagat gatantttgt cgatcaactg ttcagcacca agagagccct      420
W--> 50      ccccaacaac aactcagggt ttcacttttg gtgcttgaaa atgagtggca catgnaaaag      480
W--> 51      caagagtcnt ctttgacaaa tgtgcctgcc ganagttatc antacttact aacaagataa      540
W--> 52      tgagccaaaa catcatctgg gncatcaacc ttcatgnctt tntcaagttt atacctatna      600
W--> 53      ntnactangt cttatattnn canntgggtga ttacanttac nantaagttt agcttnaaga      660
W--> 54      aatncaagtt ttngggactc catgcctnqn cnggntttcn natccgtcgg ccagggcggn      720
W--> 55      cnggncact gntnggnagn cccanttnen cagancaeng ncccntttcc attccnggnc      780
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64 <222> LOCATION: (1)..(435)
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69      gtttatagta acttcattgc aaaactcaac aaatagattt ttggaacctt aatataataa      180
70      aattcaacag tcttctttaa ttttattctg ctcttacctt ctcataggat catatagaat      240
71      ttaaccctac aagctctcaa aaaacaatcc attattatgc tccttatcca ataaaaacaaa      300
W--> 72      accatagagt gattctcaaa atgaagattg acaaaggcaa aaagttatgc tggntcaata      360
W--> 73      gcttctttat aattntcttc atcttgccc ntccngcct tagnggtct ccattgtcaa      420
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86      tcgctcatat ggccttacag gggttgccga attagtgtga aggtaattcg gtaaattggat      120
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95 <221> NAME/KEY: misc_feature
96 <222> LOCATION: (1)..(499)
97 <223> OTHER INFORMATION: n is a, c, g, or t/u

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101      tgttttatga aggtaataca ccaattatta tggttttttg tttataaaaa tgtgaataat      180
102      tgtcaatcgt gattgcatta tctctccttt actctgtctc ttcacctttt ttaccttttt      240
103      atttgagagg aagaatccat gtagtaaaaa atgatgataa aattgttaga aaatatagtg      300
104      tcatgtaatt agagattcag attataaactt agaagacact attattttca tgaataacta      360
105      tccacgggta attatcaata ctgacatatt ttcactcaaa atattctggg tttctcatta      420
W--> 106      tatacattta aataggagct attanccatt gcaagcttgg gtttggaggc cttccgatgc      480
W--> 107      cttgttggga ttngaccca                                499
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111 <212> TYPE: DNA
112 <213> ORGANISM: Glycine max
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114 <221> NAME/KEY: misc_feature
115 <222> LOCATION: (1)..(500)
116 <223> OTHER INFORMATION: n is a, c, g, or t/u
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W--> 119      taaattgaaa atatatacgt aaganccttca tctaacagtg ctagtcgaag aatgcgtaaa      120
W--> 120      tgcagggnat ccatttccat actaaaatgg acaaaactta tatttttttt ttagcggcaa      180
W--> 121      acgttaatta ttaatttttt ttagtacaag ggatcaaacc angaccttcc ccttctttcc      240
122      atctttcttg accaccaaac caaccttata tctccacaaa acttattata tgttgttctt      300
W--> 123      cggggactat cagaattgga gtttaacctc gggcantcaa tctacataat ccttgatttn      360
W--> 124      atttngtgaa gttctaaagc cacaggcatt atttatntta ttntttctgn agtaaccnc      420
W--> 125      catatgttgg tnnataaggg tangnatnaa aatncttgg ntggtnnena tttgcncttn      480
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131 <213> ORGANISM: Glycine max
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134 <222> LOCATION: (1)..(189)
135 <223> OTHER INFORMATION: n is a, c, g, or t/u
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W--> 139      gttcattgta gcaatagcat cgttatctct tattattgct ggttcatcat natctcaatt      180
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149 <223> OTHER INFORMATION: n is a, c, g, or t

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153      taacttatta acgtttcaca ttttcaaaga cgagtctagc tatttataat tttttttcct      180
154      aaaatatatt ttttgccttc ataaatatga aaatatatta aattcgttcc taattttttt      240
155      ttcaaagcat ctttccttct cacaaaattg aaatgtatca tttttttttg ttcaaaagtt      300
156      taaataaatt tgaacctaat atgacatttt atatcggtta tacatataac tgatataaac      360
157      atcaagtttt ttatatcaat gatacctata actgatatca aatgtgacaa ttatatatat      420
158      aattaatgta aaaaagtcac aaatataatt tattttgagt caaaaaataa tatattttta      480
W--> 159      ttattttgaa gatgaaaaag gataaattta aaacatttgt gtgangatga aaaactagat      540
160      gttttttttc ctggttttaa tgcaaaacca atgctatttt atttaaattt tacctttttt      600
W--> 161      ttataattac nccacaaaaa aaccgttttg tgttacaaat ttganttaaa ttctnttggt      660
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163      atct      724
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166 <211> LENGTH: 801
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168 <213> ORGANISM: Glycine max
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171 <222> LOCATION: (1)..(801)
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176      gcacacaccc tatctttcat aaaattacta cactttttta tttttgtaat aaaaaaccta      180
177      gaaaaactca ttatgaaaca gatgatgtac ttttaacact ctgtcggcct ctctctctct      240
178      attatatatt gatttaaat tattgagaat tatatttttg ttgggtctca tttattatat      300
179      tttattaatt ggatccgggc cctctagatg cggccgcatg cataagcttg agtattctat      360
180      agtgtcacct aaatagcttg gcgtaatcat ggtcatagct gtttcctgtg tgaaattggt      420
W--> 181      atccgctcac aattccacac aacatacgag ccggaagcat aaagtgtnaa gcctggggtn      480
W--> 182      cctaattgagt gagctaactc acattaattg ccttgcgctc actgccgctt ttccagtcng      540
W--> 183      gaaacctgtc ctgccagctg cattaatgaa tngccaacc cncggggana agcngtttgc      600
W--> 184      ntatggggeg tcttnccgct tctcgcctca ntgactcget gcgctengtc nttcngntgc      660
W--> 185      cgcgaaacgt atcanncnac tcnaangnng taaatacggg tatccacna accnngggga      720
W--> 186      naaccnngga aaaaacatgt nanccaaaag gccnccaaa ggcangaaa cnttnaaaag      780
W--> 187      gccnnttgc ttgncttnt n      801
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192 <213> ORGANISM: Glycine max
193 <220> FEATURE:
194 <221> NAME/KEY: misc_feature
195 <222> LOCATION: (1)..(809)
196 <223> OTHER INFORMATION: n is a, c, g, or t
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199      ccgagctcga attcgctgac ctcgagggat ctttttatgt tggtagctac tgtaatatca      120
200      tcttgtaact ttaactttta agtcatactc ctttggact catatataag caaaagagtg      180

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203      atgataacaa gttccaatga aggacatttt agaaataacc ttatttttta ttgagatta      360
204      gtaaaattaa atgatgtgaa ctaactttct taattaatgt gaaattagtt attttttctt      420
205      atatacgagt ccaaaggagg taccaaattt cacaaatgta ctaaaatgta ttatatgctt      480
W--> 206      ctttttaatt catctttgct gcatanctac ttagctactg tgctctgata cgggacctct      540
207      agatgcggcc gcatgcataa gcttgagtat ctatagtgtc cctaaatagc ttggcgatc      600
W--> 208      atggtcatag ctgtttccng tgtgaaattg ttatccgctc acaattccac acaacatacg      660
W--> 209      anccggaagc ataaaagtgt taagccnggg gtgcctaata agtgagctaa ctcacattaa      720
W--> 210      ttgcgttgcg ctactgcc gcttcnatt cgggaaactg tcctgncanc tgcattaatg      780
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218 <221> NAME/KEY: misc_feature
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220 <223> OTHER INFORMATION: n is a, c, g, or t/u
221 <400> SEQUENCE: 11
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224      agaggggctg attttggaga aaacatcatc catggtataa agtccgttta gattccagct      180
225      attgttcaca ttcattccctt acatatgaga atatccctat aagctgaaac taacttttac      240
226      aaacaaacat gcaccgaacc attaaagtgt gacttaatat ccgggggata atgaccttaa      300
227      ttcagaaatt cacataaata actaaaagta agttgtatct tatttatgtc tggatttact      360
228      gcacaaacta aacaaaagtt tgtggattta gacataaaaa ataccaatgc tgtgtgaaaa      420
W--> 229      taagaaatgg tggatcataa gacaagtttc ttttctgttt ttttaaaatt gcagtcnaag      480
W--> 230      ccatcangag gttcatgtaa ttaaccaaac tagacgttga cttttggttt tatccttttg      540
231      tagaatagca agcaagtcac tataaatctg gccattggga cagcttagtt taactccgc      600
W--> 232      cgcaaatttg ttaaaatatt naataataat atcacctaaa atcatatttg tcanttcatt      660
W--> 233      ttgttttang ttatatcaat tattattttt taccttaant cttttataat ntcaatgatg      720
W--> 234      ggacccaaaa aattatcaa tacnttnaag onttatttat tattaattaa ncctttaatt      780
W--> 235      ataattaaaa attcnattta attttttaan      810
237 <210> SEQ ID NO: 12
238 <211> LENGTH: 777
239 <212> TYPE: DNA
240 <213> ORGANISM: Glycine max
241 <220> FEATURE:
242 <221> NAME/KEY: misc_feature
243 <222> LOCATION: (1)..(777)
244 <223> OTHER INFORMATION: n is a, c, g, or t/u
245 <400> SEQUENCE: 12
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247      catctagagg gcccgatctt ttcggttgaa gcaaaattga agtcttttgc tcatttttat      120
248      caaattcttt aatgaaaagt taattacata aaatatttta gtagaagcaa ttttacacag      180
249      ttattattta aaaaaattac acagttattc aataacaaat tacaatatat tataaggtta      240
250      taataaatat tttaaaattc atataaaaga tgacttatta ataagttgat aatgtaaatt      300
251      ttttacacta ttaaactcat ttacgtaat cttagcgaca acatactatt tttttcatga      360

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

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Seq#:5; N Pos. 92,445,494
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Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 5,6

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L:8 M:270 C: Current Application Number differs, Wrong Format

L:49 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:360
L:50 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:420
L:51 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:480
L:52 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:540
L:53 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:600
L:54 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:660
L:55 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:720
L:56 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:780
L:72 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:300
L:73 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:360
L:74 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:420
L:85 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0
L:87 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:120
L:100 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:60
L:106 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:420
L:107 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:480
L:118 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0
L:119 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:60
L:120 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:120
L:121 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:180
L:123 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:300
L:124 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:360
L:125 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:420
L:126 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:480
L:137 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0
L:138 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:60
L:139 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:120
L:159 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:480
L:161 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:600
L:162 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:660
L:174 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0
L:181 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:420
L:182 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:480
L:183 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:540
L:184 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:600
L:185 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:660
L:186 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:720
L:187 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:780
L:198 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0
L:206 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:480
L:208 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:600
L:209 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:660
L:210 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:720
L:211 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:780
L:222 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0
L:229 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:420
L:230 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:480

VERIFICATION SUMMARY

DATE: 01/14/2003

PATENT APPLICATION: US/09/954,773A

TIME: 14:52:27

Input Set : N:\paola\US09954773A.raw

Output Set: N:\CRF4\01142003\I954773A.raw

L:232 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:600
L:233 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:660
L:234 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:720